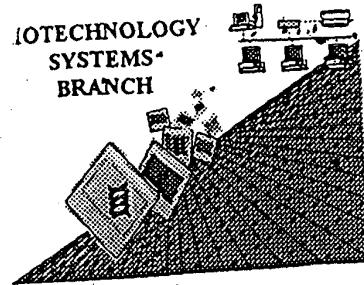0590
0920

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: ___09/865 548___

Source: ___OIPE___

Date Processed by STIC: ___09/18/2001___

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.
PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)
PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER VERSION 3.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.
Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

<u>Checker Version 3.0 can be down loaded from the USPTO website at the following address:</u>
http://www.uspto.gov/web/offices/pac/checker

ERROR DETECTED	SUGGESTED CORRECTION	SERIAL NUMBER: _O 9/ 865548_

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWA.

1 _____ Wrapped Nucleics
Wrapped Aminos

The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

2 _____ Invalid Line Length

The rules require that a line not exceed 72 characters in length. This includes white spaces.

3 _____ Misaligned Amino
Numbering

The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

4 _____ Non-ASCII

The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

5 _____ Variable Length

Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

6 _____ PatentIn 2.0
"bug"

A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

7 _____ Skipped Sequences
(OLD RULES)

Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
(i)   SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped

Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

8 _____ Skipped Sequences
(NEW RULES)

Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence.
<210> sequence id number
<400> sequence id number
000

9 _____ Use of n's or Xaa's
(NEW RULES)

Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

10 _____ Invalid <213>
Response

Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence

11 _____ Use of <220>

Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32)   (Sec. 1.823 of Sequence Rules)

12 _____ PatentIn 2.0
"bug"

Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

13 _____ Misuse of n

n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.

AMC/MH – Biotechnology Systems Branch – 08/21/2001

```
                    RAW SEQUENCE LISTING          DATE: 09/18/2001
                    PATENT APPLICATION: US/09/865,548   TIME: 15:43:18

                    Input Set : A:\PTO_VSK.txt
                    Output Set: N:\CRF3\09182001\I865548.raw

        5 <110> APPLICANT: Barnea, Eilon
        7         Beer, Ilan
        9         Ziv, Tamar
       11         Admon, Arie
       15 <120> TITLE OF INVENTION: METHOD OF IDENTIFYING PEPTIDES CAPABLE OF BINDING TO MHC
MOLECULES,
       16         PEPTIDES IDENTIFIED THEREBY AND THEIR USES
       20 <130> FILE REFERENCE: 01/22080
C--> 24 <140> CURRENT APPLICATION NUMBER: US/09/865,548
C--> 24 <141> CURRENT FILING DATE: 2001-09-04
       24 <150> PRIOR APPLICATION NUMBER: US 60/290,958
       26 <151> PRIOR FILING DATE: 2001-05-16
       30 <160> NUMBER OF SEQ ID NOS: 204
       34 <170> SOFTWARE: PatentIn version 3.0
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       40 <211> LENGTH: 9
       42 <212> TYPE: PRT
C--> 44 <213> ORGANISM: Artificial
       48 <220> FEATURE:
       50 <223> OTHER INFORMATION: synthetic peptide
       52 <400> SEQUENCE: 1
       54 Leu Leu Asp Val Pro Thr Ala Ala Val
       55 1               5
       57 <210> SEQ ID NO: 2
       59 <211> LENGTH: 10
       61 <212> TYPE: PRT
C--> 63 <213> ORGANISM: Artificial
       67 <220> FEATURE:
       69 <223> OTHER INFORMATION: synthetic peptide
       71 <400> SEQUENCE: 2
       73 Leu Leu Leu Asp Val Pro Thr Ala Ala Val
       74 1               5                   10
       76 <210> SEQ ID NO: 3
       78 <211> LENGTH: 12
       80 <212> TYPE: PRT
C--> 82 <213> ORGANISM: Artificial
       86 <220> FEATURE:
       88 <223> OTHER INFORMATION: synthetic peptide
       90 <400> SEQUENCE: 3
       92 Leu Leu Leu Asp Val Pro Thr Ala Ala Val Gln Ala
       93 1               5                   10
       95 <210> SEQ ID NO: 4
       97 <211> LENGTH: 8
       99 <212> TYPE: PRT
C--> 101 <213> ORGANISM: Artificial
      105 <220> FEATURE:
      107 <223> OTHER INFORMATION: synthetic peptide
      109 <400> SEQUENCE: 4
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Does Not Comply
Corrected Diskette Needed

Errored: Synthetic peptide is an insufficient 223 response.

FYI: "Artificial Sequence" is the preferred terminology for field 213.

The type of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

RAW SEQUENCE LISTING   DATE: 09/18/2001
PATENT APPLICATION: US/09/865,548   TIME: 15:43:18

Input Set : A:\PTO_VSK.txt
Output Set: N:\CRF3\09182001\I865548.raw

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        112 1                   5
        114 <210> SEQ ID NO: 5
        116 <211> LENGTH: 9
        118 <212> TYPE: PRT
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        126 <223> OTHER INFORMATION: synthetic peptide
        128 <400> SEQUENCE: 5
        130 Gly Leu Leu Gly Thr Leu Val Gln Leu
        131 1                   5
        133 <210> SEQ ID NO: 6
        135 <211> LENGTH: 9
        137 <212> TYPE: PRT
C--> 139 <213> ORGANISM: Artificial
        143 <220> FEATURE:
        145 <223> OTHER INFORMATION: synthetic peptide
        147 <400> SEQUENCE: 6
        149 Ala Leu Phe Gly Ala Leu Phe Leu Ala
        150 1                   5
        152 <210> SEQ ID NO: 7
        154 <211> LENGTH: 10
        156 <212> TYPE: PRT
C--> 158 <213> ORGANISM: Artificial
        162 <220> FEATURE:
        164 <223> OTHER INFORMATION: synthetic peptide
        166 <400> SEQUENCE: 7
        168 Ser Leu Leu Gly Gly Asp Val Val Ser Val
        169 1                   5                   10
        171 <210> SEQ ID NO: 8
        173 <211> LENGTH: 9
        175 <212> TYPE: PRT
C--> 177 <213> ORGANISM: Artificial
        181 <220> FEATURE:
        183 <223> OTHER INFORMATION: synthetic peptide
        185 <400> SEQUENCE: 8
        187 Asn Leu Thr Ile Ser Asp Val Ser Val
        188 1                   5
        190 <210> SEQ ID NO: 9
        192 <211> LENGTH: 9
        194 <212> TYPE: PRT
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        202 <223> OTHER INFORMATION: synthetic peptide
        204 <400> SEQUENCE: 9
        206 Ser Leu Trp Gly Gln Pro Ala Glu Ala
        207 1                   5
        209 <210> SEQ ID NO: 10
        211 <211> LENGTH: 9
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/865,548

DATE: 09/18/2001
TIME: 15:43:18

Input Set : A:\PTO_VSK.txt
Output Set: N:\CRF3\09182001\I865548.raw

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       219 <220> FEATURE:
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       223 <400> SEQUENCE: 10
       225 Ser Leu Ile Gly His Leu Gln Thr Leu
       226 1               5
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       238 <220> FEATURE:
       240 <223> OTHER INFORMATION: synthetic peptide
       242 <400> SEQUENCE: 11
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       245 1               5
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       249 <211> LENGTH: 9
       251 <212> TYPE: PRT
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       257 <220> FEATURE:
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       261 <400> SEQUENCE: 12
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       264 1               5
       266 <210> SEQ ID NO: 13
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       270 <212> TYPE: PRT
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       289 <212> TYPE: PRT
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       295 <220> FEATURE:
       297 <223> OTHER INFORMATION: synthetic peptide
       299 <400> SEQUENCE: 14
       301 Gly Leu Tyr Pro Gly Leu Ile Trp Leu
       302 1               5
       304 <210> SEQ ID NO: 15
       306 <211> LENGTH: 9
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      320 Tyr Leu Leu Pro Ala Ile Val His Ile
      321 1              5
      323 <210> SEQ ID NO: 16
      325 <211> LENGTH: 9
      327 <212> TYPE: PRT
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      337 <400> SEQUENCE: 16
      339 Ala Leu Ser Asp His His Ile Tyr Leu
      340 1              5
      342 <210> SEQ ID NO: 17
      344 <211> LENGTH: 9
      346 <212> TYPE: PRT
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      352 <220> FEATURE:
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      356 <400> SEQUENCE: 17
      358 Ile Leu Asp Gln Lys Ile Asn Glu Val
      359 1              5
      361 <210> SEQ ID NO: 18
      363 <211> LENGTH: 9
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      371 <220> FEATURE:
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      377 Ile Leu Asp Lys Lys Val Glu Lys Val
      378 1              5
      380 <210> SEQ ID NO: 19
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      384 <212> TYPE: PRT
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      390 <220> FEATURE:
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      397 1              5                  10
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C--> 405 <213> ORGANISM: Artificial
      409 <220> FEATURE:
      411 <223> OTHER INFORMATION: synthetic peptide
      413 <400> SEQUENCE: 20
      415 Gly Val Tyr Asp Gly Glu Glu His Ser Val
      416 1              5                  10
      418 <210> SEQ ID NO: 21
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       422 <212> TYPE: PRT
C--> 424 <213> ORGANISM: Artificial
       428 <220> FEATURE:
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       432 <400> SEQUENCE: 21
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       435 1               5                   10
       437 <210> SEQ ID NO: 22
       439 <211> LENGTH: 9
       441 <212> TYPE: PRT
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       447 <220> FEATURE:
       449 <223> OTHER INFORMATION: synthetic peptide
       451 <400> SEQUENCE: 22
       453 Thr Leu Trp Val Asp Pro Tyr Glu Val
       454 1               5
       456 <210> SEQ ID NO: 23
       458 <211> LENGTH: 10
       460 <212> TYPE: PRT
C--> 462 <213> ORGANISM: Artificial
       466 <220> FEATURE:
       468 <223> OTHER INFORMATION: synthetic peptide
       470 <400> SEQUENCE: 23
       472 Phe Leu Phe Asp Gly Ser Pro Thr Tyr Val
       473 1               5                   10
       475 <210> SEQ ID NO: 24
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       479 <212> TYPE: PRT
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       485 <220> FEATURE:
       487 <223> OTHER INFORMATION: synthetic peptide
       489 <400> SEQUENCE: 24
       491 Phe Leu Phe Asp Gly Ser Pro Thr Tyr Val Leu
       492 1               5                   10
       494 <210> SEQ ID NO: 25
       496 <211> LENGTH: 12
       498 <212> TYPE: PRT
C--> 500 <213> ORGANISM: Artificial
       504 <220> FEATURE:
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       508 <400> SEQUENCE: 25
       510 Ala Leu Trp Asp Ile Glu Thr Gly Gln Gln Thr Val
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       513 <210> SEQ ID NO: 26
       515 <211> LENGTH: 9
       517 <212> TYPE: PRT
C--> 519 <213> ORGANISM: Artificial
       523 <220> FEATURE:
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VERIFICATION SUMMARY   DATE: 09/18/2001
PATENT APPLICATION: US/09/865,548   TIME: 15:43:19

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Output Set: N:\CRF3\09182001\I865548.raw

L:24 M:270 C: Current Application Number differs, Replaced Current Application No
L:24 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:44 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:1
L:63 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:2
L:82 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
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VERIFICATION SUMMARY   DATE: 09/18/2001
PATENT APPLICATION: US/09/865,548   TIME: 15:43:19

Input Set : **A:\PTO_VSK.txt**
Output Set: **N:\CRF3\09182001\I865548.raw**

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L:973 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:50